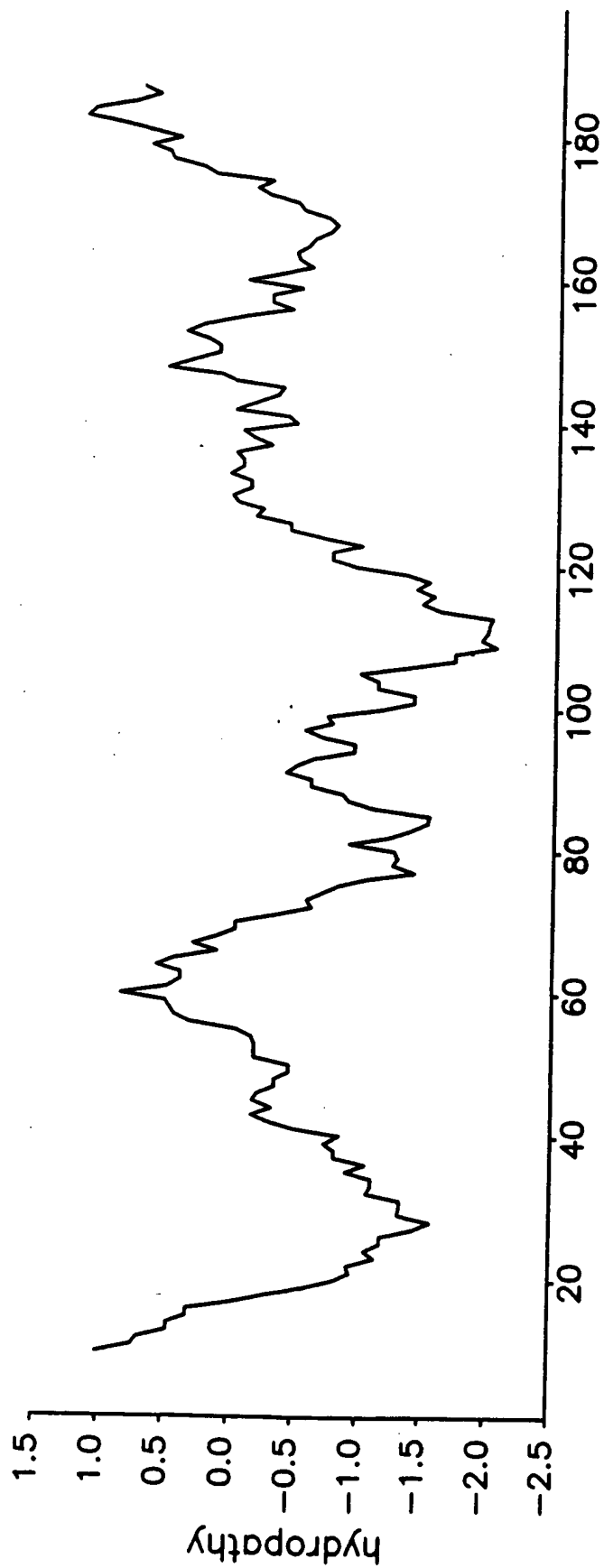


home/ruby/va/Molbio/nico/DNA6/p1.DNA62377 (length: 197)
kyte (hydropathy); window: 20



IL 17 biology #2

DNA 62377

```
> /nico Unix Files/DNA6/ss.DNA62377
> 0 Sites [No Sites]
> length: 1047 bp (circular)
```

```
> /usr/seqdb2/sst/DNA/Dnaseqs.full/ss.DNA62377 (1250 bases)
[DNA62377], sheldens
```

>insert starts here

```

1 GCCAGGTGTG CAGGCCGCTC CAAGCCACAGC CTGCCCCCGCT GCCGCCACCA TGACGCTCCT CCGGGCCTC CTGTTTCTGA CCTGGCTGCA CACATGCCTG
  CGGTCCACAC GTCCGGCGAG GTTCGGGTG GACGGGGCGA CGCGGTGCT ACTGCGAGGA GGGGCGGAG GACAAAGACT GGACCGACGT GTGTACGGAC
1
      M T L L P G L L F L T W L H T C L
      ^MET

101 GCCCACCATG ACCCTCCCT CAGGGGGCAC CCCCACAGTC ACGGTACCC ACACGTCTAC TCGGCTGAGG AACTGCCCT CGGCCAGGCC CCCCCACACC
    CGGTGTGTAC TGGGGAGGA GTCCCCCGTG GGGGTGTCTAG TGCCATGGG TGTCACGATG AGCCGACTCC TTGACGGGA GCCGGTCCGG GGGGTGTGG
18 A H H D P S L R G H P H S H G T P H C Y S A E E L P L G Q A P P H L

201 TGCTGGCTCG AGGTGCCAAG TGGGGGCAGG CTTTGCCTGT AGCCTGGTG TCCAGCCTGG AGGCAGCAAG CCACAGGGGG AGGCACGAGA GGCCCTCAGC
    ACGACCGAGC TCCACGGTTC ACCCCCGTCC GAAACGGACA TCGGGACCAC AGGTGGAGC TCCGTCTGTC GGTGTCCCC TCCGTGCTCT CCGGGAGTCG
52 L A R G A K W G Q A L P V A L V S S L E A A S H R G R H E R P S A

301 TAGCACCCAG TGCCCGGTGC TGCGGCCGGA GGAGGTGTTG GAGGCAGACA CCACACAGCG CTCCATCTCA CCCTGGAGAT ACCGTGTGGA CACGGATGAG
    ATGCTGGTTC ACGGGCCACG ACGCCGGCCT CCTCCACAAC CTCGCTGTGT GGTGGTCTG GAGGTAGAT GGGACCTCTA TGGCACACCT GTGCCTACTC
85 T T Q C P V L R P E E V L E A D T H Q R S I S P W R Y R V D T D E

401 GACCGCTATC CACAGAAGCT GGCCTTCGCC GAGTGCCTGT GCAGAGGCTG TATCGATGCA CGGACGGGCC GCGAGACAGC TGCCTCAAC TCCGTGCGGC
    CTGGCGATAG GTGTCTTCGA CCGGAAGCGG CTCACGGACA CGTCTCCGAC ATAGCTACGT GCCTGCCCGG CGCTCTGTG ACGCAGATTG AGGCACGCCG
118 D R Y P Q K L A F A E C L C R G C I D A R T G R E T A A L N S V R L

501 TGCTCCAGAG CTGTGTTGTTG CTGCGCCGCC GGCCCTGCTC CCGGACGGC TCGGGGCTCC CCACACCTGG GGCCTTTGCC TTCCACACCG AGTTCATCCA
    ACGAGGTCTC GGACGACCAC GACGGCGCGG CCGGGACGAG GCGCTGCCG AGCCCGAGG GGTGTGGACC CCGGAAACGG AAGGTGTGGC TCAAGTAGGT
152 L Q S L L V L R R R P C S R D G S G L P T P G A F A F H T E F I H

601 CGTCCCCGTC GGCTGCACCT GCGTGTGCTC CCGTTTCAGT TGACCGCGCA GGGCGTGGG CCCCTAGACT GGACACGTGT GCTCCCCAGA GGGCACCCCC
    GCAGGGGCAG CCGACGTGGA CGCACGACGG GGCAAGTCAC ACTGGCGGCT CCGGCACCCC GGGGATCTGA CCTGTGCACA CGAGGGGTCT CCCGTGGGGG
185 V P V G C T C V L P R S V O

701 TATTATATG TATTATTGT TATTATATG CCTCCCCCAA CACTACCTT GGGTCTGGG CATTCCCCGT GTCTGGAGGA CAGCCCCCA CTGTTCTCT
    ATAAATACAC ATAAATAACA ATAAATATAC GGAGGGGGTT GTGATGGAA CCCCAGACCC GTAAGGGCA CAGACCTCCT GTCGGGGGT GACAAGAGGA
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801 CATCTCCAGC CTCAGTAGTT GGGGGTAGAA GGAGCTCAGC ACCTCTTCCA GCCCTTAAAG CTGCAGAAA GGTGTCACAC GGCTGCCTGT ACCTTGGCTC
 GTAGAGGTCG GAGTCATCAA CCCCATCTT CCTCGAGTCG TGGAGAAGGT CCGGAATTTC GACGCTTTT CCACAGTGT CCGACGGACA TGGAAACCGAG
 901 CCTGTCTCTGC TCCCGGCTTC CCTTACCCTA TCACTGGCCT CAGGCCCCGC AGGCTGCCTC TTCCCAACCT CCTTGGAAAGT ACCCCTGTTT CTTAAACAAT
 GGACAGGACG AGGCCGAAG GGAATGGAT AGTGACCGGA GTCCGGGGCG TCCGACGGAG AAGGGTTGGA GGAACCTTCA TGGGGACAAA GAATTTGTTA
 1001 TATTTAAGTG TAGGTGATT ATTAACTGA TGAACACATC CCCAAA
 ATAAATTCAC ATGCACATAA TAATTGACT ACTTGTGTAG GGGTTTT